

Day 4B

Bootstrap

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1. Introduction

- The bootstrap is a method for obtaining properties of statistics through resampling.
- There are many ways to bootstrap.
- There are many uses of the bootstrap.
- The most common uses of the bootstrap in econometrics are
 - ▶ to obtain standard errors of estimates.
- Occasionally use a more advanced bootstrap to potentially enable better finite sample inference
 - ▶ confidence intervals with better coverage
 - ▶ tests with true size closer to nominal size.

Summary

- 1 Introduction
- 2 Bootstrap (without asymptotic refinement)
- 3 Bootstrap in General
- 4 Bootstrap with asymptotic refinement
- 5 Wild Bootstrap
- 6 Stata commands

2. Bootstrap pairs estimate of standard error

- The most common bootstrap is the pairs bootstrap
 - ▶ views the sample as the $\{(y_1, \mathbf{x}_1), \dots, (y_N, \mathbf{x}_N)\}$ as the population
 - ▶ assumes that (y_i, \mathbf{x}_i) are i.i.d.
 - ▶ obtains B random samples from this population by resampling with replacement
 - ★ e.g. in bootstrap resample 1 observation may appear once, observation 2 not at all, observation 2 times,
- This yields B estimates $\hat{\theta}_1, \dots, \hat{\theta}_B$.
 - ▶ so estimate $\text{Var}[\hat{\theta}]$ using the usual variance of the B estimates.
- For scalar θ we have

$$\widehat{\text{V}}[\hat{\theta}] = \frac{1}{B-1} \sum_{b=1}^B (\hat{\theta}_b - \bar{\hat{\theta}})^2, \quad \text{where } \bar{\hat{\theta}} = \frac{1}{B} \sum_{b=1}^B \hat{\theta}_b.$$

- ▶ Square root of this is called a bootstrap standard error.

Regression application

- Data: Doctor visits (count) and chronic conditions. $N = 50$.

```
. * Summarize and Poisson with robust se's
. summarize
```

variable	Obs	Mean	Std. Dev.	Min	Max
docvis	50	4.12	7.82106	0	43
age	50	4.162	1.160382	2.6	6.2
chronic	50	.28	.4535574	0	1

```
. poisson docvis chronic, nolog vce(robust)
```

```
Poisson regression                                Number of obs   =           50
                                                    wald chi2(1)    =           3.64
                                                    Prob > chi2     =          0.0565
Log pseudolikelihood = -238.75384                Pseudo R2      =          0.0917
```

docvis	Coef.	Robust Std. Err.	z	P> z	[95% Conf. Interval]	
chronic	.9833014	.5154894	1.91	0.056	-.0270391	1.993642
_cons	1.031602	.3446734	2.99	0.003	.3560541	1.707149

Bootstrap standard errors after Poisson regression

- Use option `vce(boot)`
 - ▶ Set the seed!
 - ▶ Set the number of bootstrap repetitions!

```
. * Compute bootstrap standard errors using option vce(bootstrap) to
. poisson docvis chronic, vce(boot, reps(400) seed(10101) nodots)
```

```
Poisson regression                                Number of obs   =          50
                                                Replications    =          400
                                                Wald chi2(1)    =           3.33
                                                Prob > chi2     =          0.0679
Log likelihood = -238.75384                       Pseudo R2       =          0.0917
```

docvis	Observed Coef.	Bootstrap Std. Err.	z	P> z	Normal-based [95% Conf. Interval]	
chronic	.9833014	.5386575	1.83	0.068	-.0724478	2.039051
_cons	1.031602	.3536507	2.92	0.004	.338459	1.724744

- Bootstrap se = 0.539 versus White robust se = 0.515.

Results vary with seed and number of reps

```

. * Bootstrap standard errors for different reps and seeds
. quietly poisson docvis chronic, vce(boot, reps(50) seed(10101))

. estimates store boot50

. quietly poisson docvis chronic, vce(boot, reps(50) seed(20202))

. estimates store boot50diff

. quietly poisson docvis chronic, vce(boot, reps(2000) seed(10101))

. estimates store boot2000

. quietly poisson docvis chronic, vce(robust)

. estimates store robust

. estimates table boot50 boot50diff boot2000 robust, b(%8.5f) se(%8.5f)

```

variable	boot50	boot50~f	boot2000	robust
chronic	0.98330	0.98330	0.98330	0.98330
	0.45444	0.59923	0.54178	0.51549
_cons	1.03160	1.03160	1.03160	1.03160
	0.37131	0.37533	0.36414	0.34467

Legend: b/se

Leading uses of bootstrap standard errors

- Sequential two-step m-estimator
 - ▶ First step gives $\hat{\alpha}$ used to create a regressor $z(\hat{\alpha})$
 - ▶ Second step regresses y on x and $z(\hat{\alpha})$
 - ▶ Do a paired bootstrap resampling (x, y, z)
 - ▶ e.g. Heckman two-step estimator.
- 2SLS estimator with heteroskedastic errors (if no White option)
 - ▶ Paired bootstrap gives heteroskedastic robust standard errors.
- Functions of other estimates e.g. $\hat{\theta} = \hat{\alpha} \times \hat{\beta}$
 - ▶ replaces delta method
 - ▶ Clustered data with many small clusters, such as short panels.
 - ★ Then resample the clusters.
 - ★ But be careful if model includes cluster-specific fixed effects.
- For these in Stata need to use prefix command `bootstrap`:

3. The Bootstrap in General: Bootstrap algorithm

- A general bootstrap algorithm is as follows:
 - ▶ **1.** Given data $\mathbf{w}_1, \dots, \mathbf{w}_N$
 - ★ draw a bootstrap sample of size N (see below)
 - ★ denote this new sample $\mathbf{w}_1^*, \dots, \mathbf{w}_N^*$.
 - ▶ **2.** Calculate an appropriate statistic using the bootstrap sample. Examples include:
 - ★ (a) estimate $\hat{\theta}^*$ of θ ;
 - ★ (b) standard error $s_{\hat{\theta}}^*$ of estimate $\hat{\theta}^*$
 - ★ (c) t -statistic $t^* = (\hat{\theta}^* - \hat{\theta}) / s_{\hat{\theta}}^*$ centered at $\hat{\theta}$.
 - ▶ **3.** Repeat steps 1-2 B independent times.
 - ★ Gives B bootstrap replications of $\hat{\theta}_1^*, \dots, \hat{\theta}_B^*$ or t_1^*, \dots, t_B^* or
 - ▶ **4.** Use these B bootstrap replications to obtain a bootstrapped version of the statistic (see below).

Implementation

- Number of bootstraps: B high is best but increases computer time.
 - ▶ CT use 400 for se's and 999 for tests and confidence intervals.
 - ▶ Defaults are often too low. And set the seed!
- Various resampling methods
 - ▶ 1. Paired (or nonparametric or empirical dist. func.) is most common
 - ★ $\mathbf{w}_1^*, \dots, \mathbf{w}_N^*$ obtained by sampling with replacement from $\mathbf{w}_1, \dots, \mathbf{w}_N$.
 - ▶ 2. Parametric bootstrap for fully parametric models.
 - ★ Suppose $y|\mathbf{x} \sim F(\mathbf{x}, \theta_0)$ and generate y_i^* by draws from $F(\mathbf{x}_i, \hat{\theta})$
 - ▶ 3. Residual bootstrap for regression with additive errors
 - ★ Resample fitted residuals $\hat{u}_1, \dots, \hat{u}_N$ to get $(\hat{u}_1^*, \dots, \hat{u}_N^*)$ and form new $(y_1^*, \mathbf{x}_1), \dots, (y_N^*, \mathbf{x}_N)$.
- Need to resample over i.i.d. observations
 - ▶ resample over clusters if data are clustered
 - ★ But be careful if model includes cluster-specific fixed effects.
 - ▶ resample over moving blocks if data are serially correlated.

Bootstrap failure

- The bootstrap always provides estimates even when it makes no sense
 - ▶ e.g. can always get bootstrap standard errors for the mean of a Cauchy sample, even though the mean of the Cauchy does not exist.
- The following are cases where standard bootstraps fail
 - ▶ so need to adjust standard bootstraps.
- GMM (and empirical likelihood) in over-identified models
 - ▶ For overidentified models need to recenter or use empirical likelihood.
- Nonparametric Regression:
 - ▶ Nonparametric density and regression estimators converge at rate less than \sqrt{N} and are asymptotically biased.
 - ▶ This complicates inference such as confidence intervals.
- Non-Smooth Estimators: e.g. LAD.

Jackknife

- The jackknife uses a leave-one-out resampling scheme.
- The jackknife estimate of the variance of an estimator $\hat{\theta}$ is

$$\widehat{V}[\hat{\theta}] = \frac{N-1}{N} \sum_{i=1}^N (\hat{\theta}_{(-i)} - \bar{\hat{\theta}})^2, \quad \text{where } N^{-1} \sum_i \hat{\theta}_{(-i)}.$$

- ▶ where $\hat{\theta}_{(-i)}$ is $\hat{\theta}$ obtained from the sample with observation i omitted.
- The jackknife is a “rough and ready” method for bias reduction in many situations, but not the ideal method in any.
 - ▶ it can be viewed as a linear approximation of the bootstrap (Efron and Tibsharani (1993, p.146)).
 - ▶ it requires less computation than the bootstrap in small samples, as then $N < B$ is likely
 - ▶ but it is outperformed by the bootstrap as $B \rightarrow \infty$.
- E.g. `poisson docvis chronic, vce(jackknife)`

4. Bootstrap with asymptotic refinement

- The simplest bootstraps are no better than usual asymptotic theory
 - ▶ advantage is easy to implement, e.g. standard errors.
- More complicated bootstraps provide asymptotic refinement
 - ▶ this may provide a better finite-sample approximation.
- Conventional asymptotic tests (such as Wald test).
 - ▶ α = nominal size for a test, e.g. $\alpha = 0.05$.
 - ▶ Actual size = $\alpha + O(N^{-1/2})$.
- Tests with asymptotic refinement
 - ▶ Actual size = $\alpha + O(N^{-1})$.
 - ▶ asymptotic bias of size $O(N^{-1}) < O(N^{-1/2})$ is smaller asymptotically.
 - ▶ But need simulation studies to confirm finite sample gains.
 - ★ e.g. if $N = 100$ then $100/N = O(N^{-1}) > 5/\sqrt{N} = O(N^{-1/2})$.

Asymptotically pivotal statistic and studentized t-statistic

- Econometricians rarely use asymptotic refinement.
- Asymptotic refinement bootstraps an asymptotically pivotal statistic
 - ▶ this means limit distribution does not depend on unknown parameters.
- An estimator $\hat{\theta} - \theta_0 \stackrel{a}{\sim} \mathcal{N}[0, \sigma_{\hat{\theta}}^2]$ is not asymptotically pivotal
 - ▶ since $\sigma_{\hat{\theta}}^2$ is an unknown parameter.
- But the studentized t -statistic is asymptotically pivotal
 - ▶ since $t = (\hat{\theta} - \theta_0) / s_{\hat{\theta}} \stackrel{a}{\sim} \mathcal{N}[0, 1]$ has no unknown parameters.
- So bootstrap Wald test statistic to get tests and confidence intervals with asymptotic refinement.
- Formally this is an empirical way of implementing an Edgeworth expansion
 - ▶ a higher order expansion than the usual one used for asymptotic theory
 - ▶ analogous to going out extra terms in a Taylor series expansion.

Percentile-t bootstrap

- Bootstrap $t = (\hat{\theta} - \theta_0) / s_{\hat{\theta}} \stackrel{a}{\sim} \mathcal{N}[0, 1]$
 - ▶ by recomputing $t_b^* = (\hat{\theta}_b - \hat{\theta}) / s_{\hat{\theta}_b}$ where $\hat{\theta} =$ original sample estimate.

```
. * Percentile-t for a single coefficient: Bootstrap the t statistic
. use bootdata.dta, clear

. quietly poisson docvis chronic, vce(robust)

. local theta = _b[chronic]

. local setheta = _se[chronic]

. bootstrap tstar=((_b[chronic]-`theta')/_se[chronic]), seed(10101) ///
> reps(999) nodots saving(percentilet, replace): poisson docvis chronic, ///
> vce(robust)
(note: file percentilet.dta not found)
```

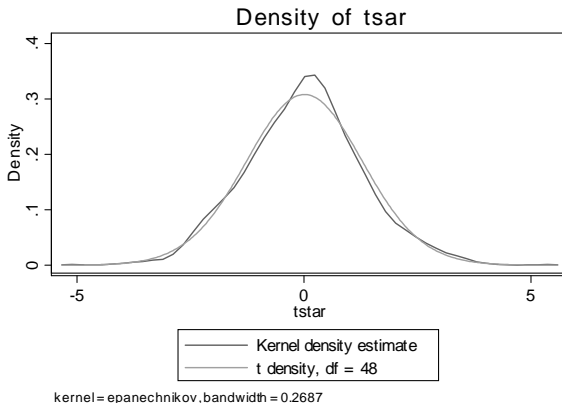
```
Bootstrap results                                Number of obs    =        50
                                                Replications     =       999
```

```
command: poisson docvis chronic, vce(robust)
tstar:   (_b[chronic]-.9833014421442415)/_se[chronic]
```

	Observed Coef.	Bootstrap Std. Err.	z	P> z	Normal-based [95% Conf. Interval]	
tstar	0	1.288018	0.00	1.000	-2.52447	2.52447

Percentile-t bootstrap (continued)

- The 999 values of t_{star} ($= t_b^* = (\hat{\theta}_b - \hat{\theta}) / s_{\hat{\theta}_b}$) trace the bootstrap estimated density of the t-statistic .
- The plot is of the kernel density estimate and $T(48)$



Percentile-t bootstrap (continued)

- The test and confidence interval critical t -values are the 2.5 and 97.5 percentiles of t^*
- For a symmetric two-sided test of $H_0 : \theta = 0$ find the proportion of times that $|t^*| > |t|$

```
. * Percentile-t critical values, 95% confidence interval
. * and p-value for symmetric two-sided wald test of H0: theta = 0
. use percentile, clear
(bootstrap: poisson)

. centile tstar, c(2.5, 97.5)
```

variable	Obs	Percentile	Centile	— Binom. Interp. — [95% Conf. Interval]	
tstar	999	2.5	-2.414266	-2.566386	-2.296212
		97.5	2.661827	2.457073	3.011566

```
. display "95% conf. int. = (" .983-r(c_1)*0.515 ", " .983+r(c_2)*0.515 ")"
95% conf. int. = (2.226347, 2.3538408)

. * Percentile-t p-value for symmetric two-sided wald test of H0: theta = 0
. quietly count if abs(tstatistic) < abs(tstar)

. display "tstatistic = " tstatistic " and p-value = " r(N)/(_N+1)
tstatistic = 1.9075106 and p-value = .146
```

BC and BCa confidence interval

- The BC (bias-corrected) and BCa methods also provide asymptotic refinement.
- (N) is observed coefficient $\pm 1.96 \times$ bootstrap s.e.
- (P) is 2.5 to 97.5 percentile of the bootstrap estimates $\hat{\beta}_1^*, \dots, \hat{\beta}_B^*$.
- (BC) and (BCa) have asymptotic refinement.

```
. * Bootstrap confidence intervals: normal-based, percentile, BC, and BCa
. quietly poisson docvis chronic, vce(boot, reps(999) seed(10101) bca)
```

```
. estat bootstrap, all
```

```
Poisson regression                Number of obs    =          50
                                Replications      =          999
```

docvis	Observed Coef.	Bias	Bootstrap Std. Err.	[95% Conf. Interval]		
chronic	.98330144	.0132307	.54137854	-.077781	2.044384	(N)
				-.0139438	2.061742	(P)
				-.079079	2.019438	(BC)
				.0295944	2.08349	(BCa)
_cons	1.0316016	-.0769223	.35685342	.3321817	1.731021	(N)
				.186586	1.582409	(P)
				.268264	1.641356	(BC)
				.386773	1.771351	(BCa)

```
(N) normal confidence interval
```

```
(P) percentile confidence interval
```

```
(BC) bias-corrected confidence interval
```

```
(BCa) bias-corrected and accelerated confidence interval
```

5. Wild bootstrap

- The wild bootstrap is specific to linear regression.
- It conditions on the sample value of the \mathbf{x}' s.
- Only y is resampled. \mathbf{x} is not resampled.
- In original sample do OLS of y_i on \mathbf{x}_i and get residual $\hat{u}_i = y_i - \mathbf{x}_i' \hat{\beta}$.
- In the b^{th} resample

$$\blacktriangleright \text{ set } y_{i,b} = \begin{cases} \mathbf{x}_i' \hat{\beta} + \hat{u}_i & \text{with probability 0.5} \\ \mathbf{x}_i' \hat{\beta} - \hat{u}_i & \text{with probability 0.5} \end{cases}$$

- ▶ do OLS regression using sample $(y_{1,b}, \mathbf{x}_1), \dots, (y_{N,b}, \mathbf{x}_N)$ gives $t_b^* = (\hat{\beta}_b - \hat{\beta}) / s_{\hat{\beta}_b}$.
 - ▶ seems "wild" as $y_{i,b}$ can only take one of two values
 - ▶ but we are doing this for N observation so possibly as many as 2^N distinct samples.
- Gives an asymptotic refinement for OLS with heteroskedastic errors.

Wild bootstrap (continued)

- Not used much in practice for independent observations.
 - ▶ usually if N is so low that we need this bootstrap as asymptotic theory has kicked in then things are statistically insignificant.
- But with clustered data things can be highly statistically significant
 - ▶ yet tests have poor size if number of clusters G is small
 - ▶ Cameron, Gelbach and Miller proposed a cluster version of the Wild bootstrap where resample over clusters
 - ▶ i.e. $\mathbf{y}_{g,b} = \mathbf{X}'_g \hat{\boldsymbol{\beta}} + \hat{\mathbf{u}}_g$ or $\mathbf{y}_{g,b} = \mathbf{X}'_g \hat{\boldsymbol{\beta}} - \hat{\mathbf{u}}_g$ in cluster g
 - ▶ and then use the percentile-t method as before
- In theory one could instead use a simpler pairs cluster bootstrap
 - ▶ where resample clusters $(\mathbf{y}_g, \mathbf{X}_g)$ with replacement
 - ▶ but this worked poorly in Monte Carlos.

6. Stata commands

- Most commands have option `vce(bootstrap)` and `vce(jackknife)`
- For more complicated bootstraps write a program and use `bootstrap`:
- For replicability set the seed!!
- For published work the more bootstraps the better as the seed becomes less important.

7. References

- A. Colin Cameron and Pravin K. Trivedi (2005), *Microeconometrics: Methods and Applications (MMA)*, chapter 11, Cambridge Univ. Press.
- A. Colin Cameron and Pravin K. Trivedi (2009), *Microeconometrics using Stata (MUS)*, chapter 13, Stata Press.